

SEQUENCE LISTING

<110>	Braun, Klaus Braun, Isabell Debus, Jürgen Pipkorn, Rüdiger Waldeck, Waldemar	
<120>	CONJUGATE FOR TREATING PROKARYOTIC INFECTIONS	
<130>	4121-170	
<140> <141>	10/501,962 2004-07-19	
<150> <151>	PCT/DE03/00124 2003-01-17	
<150> <151>	DE 10201862.6 2002-01-18	
<160>	31	
<170>	PatentIn version 3.3	
<210> <211> <212> <213>	1 15 DNA Artificial Sequence	
<220> <223>	Synthetic Construct	
<400> 1 attgttagat ttcat		15
<210><211><211><212><213>	2 14 DNA Artificial Sequence	
<220> <223>	Synthetic Construct	
<400> tcttgt	2 tcaa tcat	14
<210><211><211><212><213>	3 600 DNA Artificial Sequence	
<220> <223>	Synthetic Construct	
<400> ttctca	3 etgtt tgacagetta teategataa getttaatge ggtagtttat eacagttaaa	60
ttgcta	acgc agtcaggcac cgtgtatgaa atctaacaat gcgctcatcg tcatcctcgg	120
caccgt	cace etggatgetg taggeatagg ettggttatg eeggtaetge egggeetett	180

170 Sequence Listing.ST25.txt

240

300

360

420

480

540

600

```
gcgggatatc gtccattccg acagcatcgc cagtcactat ggcgtgctgc tagcgctata
tgcqttgatg caatttctat gcgcacccgt tctcggagca ctgtccgacc gctttggccg
ccgcccagtc ctgctcgctt cgctacttgg agccactatc gactacgcga tcatggcgac
cacacccgtc ctgtggatcc tctacgccgg acgcatcgtg gccggcatca ccggcgccac
aggtgcggtt gctggcgcct atatcgccga catcaccgat ggggaagatc gggctcgcca
cttcgggctc atgagcgctt gtttcggcgt gggtatggtg gcaggccccg tggccggggg
actgttgggc gccatctcct tgcatgcacc attccttgcg gcggcggtgc tcaacggcct
<210> 4 .
<211> 109
<212> PRT
<213> Bacteriophage P1
<400> 4
Met Leu Asp Thr Gln Glu Leu Ala Pro Val Ala Ile Ala Leu Leu Leu
Ser Val Ile Gly Gly Ile Gly Thr Phe Leu Met Asp Val Arg Asp Gly
Arg Gln Ser Gly Asn Leu Leu Gly Leu Val Thr Glu Ile Phe Val Ala
                           40
Val Thr Ala Gly Ala Val Ala Tyr Leu Leu Gly Gln His Glu Gly Trp
Glu Leu Ser Ile Thr Tyr Leu Met Val Thr Ile Ala Ser Asn Asn Gly
His Glu Val Ile Ser Gly Met Lys Arg Val Asn Ile Asp Ser Ile Leu
Asn Val Leu Thr Ser Leu Val Lys Lys Gly Gly Gly Lys
<210> 5
<211> 68
<212> PRT
<213> Bacteriophage H19B
<400> 5
Met Glu Lys Ile Thr Thr Gly Val Ser Tyr Thr Thr Ser Ala Val Gly
```

Thr Gly Tyr Trp Leu Leu Gln Leu Leu Asp Lys Val Ser Pro Ser Gln

20

Trp Val Ala Ile Gly Val Leu Gly Ser Leu Leu Phe Gly Leu Leu Thr 40 Tyr Leu Thr Asn Leu Tyr Phe Lys Ile Arg Glu Asp Arg Arg Lys Ala Val Arg Gly Glu <210> 6 <211> 96 <212> PRT <213> Bacteriophage A118 <400> 6 Met Ile Glu Met Glu Phe Gly Lys Glu Leu Leu Val Tyr Met Thr Phe Leu Val Val Val Thr Pro Val Phe Val Gln Ala Ile Lys Lys Thr Glu Leu Val Pro Ser Lys Trp Leu Pro Thr Val Ser Ile Leu Ile Gly Ala 40 Ile Leu Gly Ala Leu Ala Thr Phe Leu Asp Gly Ser Gly Ser Leu Ala 55 Thr Met Ile Trp Ala Gly Ala Leu Ala Gly Ala Gly Gly Thr Gly Leu Phe Glu Gln Phe Thr Asn Arg Ser Lys Lys Tyr Gly Glu Asp Asp Lys <210> 7 <211> 143 <212> PRT <213> Lactobacillus casei bacteriophage A2 <400> 7 Met Lys Ile Asn Trp Lys Val Ala Val Leu Ser Val Lys Phe Trp Leu 10 Ala Leu Val Pro Ala Ala Leu Leu Val Val Gln Thr Ala Ala Ala Val

Val Ile Asn Ala Val Phe Ala Leu Leu Thr Ile Val Gly Val Ala Val

Phe Gly Tyr Asn Trp Asp Phe Ala Asn Leu Gly Lys Glu Leu Thr Ala

50 55 Asp Pro Thr Thr Glu Gly Val Ser Asp Ser Gln Gln Ala Leu Ala Tyr Pro Ala Leu Ile Thr Thr Lys Ala Ala Lys Ile Lys Ser Leu Glu Asp Gln Ile Lys Ala Leu Gln Ala Asp Lys Ala Ala Asp Gln Ala Thr Ser 105 Ala Ala Ser Glu Val Val Pro Glu Thr Ser Ser Ala Ala Pro Ala Glu 120 Ser Ala Pro Glu Ser Val Ala Pro Val Ala Ser Glu Glu Val Lys 135 <210> 8 <211> 142 <212> PRT <213> Lactobacillus bacteriophage phig le <400> 8 Met Asp Ile Ile Thr Ser Leu Asn Leu Ala Thr Ala Gly Glu Leu Ala Leu Ile Ser Phe Phe Ile Gly Val Ile Val Gln Ala Ile Lys Lys Thr Gly Lys Val Lys Asn Thr Tyr Leu Pro Phe Ile Ser Met Gly Ile Gly Ile Leu Ala Gly Leu Ala Ala Val Val Thr Lys Asp Thr Asn Tyr Leu Asn Gly Ala Val Ala Gly Leu Ile Val Gly Ala Ala Thr Ser Gly Leu Thr Asp Gly Leu Ser Val Gly Thr Ser Ala Val Thr Thr Ala Lys Ala Thr Lys Asp Ala Ala Lys Thr Ala Ala Ile Thr Gln Ala Val Leu Asn Ser Ile Asn Thr Thr Lys Ser Ser Asp Thr Thr Gln Val Ala Asn 115 120

Thr Ser Asn Thr Glu Gly Gly Ser Thr Ser Glu Thr Gln Lys

```
<400> 9
Met Thr Leu Ile Asp Trp Phe Asn Leu Ile Val Ala Ile Gly Thr Ile
Ala Leu Ala Val Val Ala Ser Val Tyr Val His Leu Lys Ala Lys Ile
                             25
Asp Thr Lys Thr Ala Ala Gly Lys Ala Phe Asp Leu Val Gly Lys Leu
Ala Val Trp Ala Val Asn Glu Ala Glu His Ser Gln Asp Gly Gly Ala
Ala Lys Arg Glu Phe Ala Ala Lys Leu Ile Ser Asp Gln Leu Lys Ala
Lys Gly Ile Thr Gly Ile Asp Glu Lys Met Val Tyr Gly Ala Val Glu
Thr Ala Trp Lys Glu Ala Ile Glu Asn Val Lys
           100
<210> 10
<211> 44
<212> PRT
<213> Lactococcus phage c2
<400> 10
Met Ile Glu Thr Leu Arg Ala Ile Gly Leu Val Val Phe Met Gln Leu
Leu Ser Leu Ala Leu Glu Phe Ile Asp Thr Gly Thr Leu Lys Pro Ser
Val Arg Lys Arg Ile Ala Val Glu Leu Met Val Leu
                         40
<210> 11
<211> 74
<212>
      PRT
<213> bacteriophage phi AM2
<400> 11
Met Phe Phe Asn Asn Lys Phe Tyr Asn Val Ile Lys Trp Ala Val Leu
```

<210> 9

10 15 5 1 Thr Ala Leu Pro Ala Leu Ser Val Phe Ile Gly Val Ile Gly Lys Ala 25 Tyr Gly Trp Gly Gly Thr Asp Leu Ala Ile Ile Thr Leu Asn Ala Phe 40 Thr Val Phe Leu Gly Thr Leu Ala Gly Val Ser Ala Val Lys Tyr Asn Ser Gln Pro Asn Asp Thr Lys Glu Asn Lys 70 <210> 12 <211> 88 <212> PRT <213> Bacteriophage Tuc2009 <400> 12 Met Asn Gln Ile Asn Trp Lys Leu Arg Leu Lys Ser Lys Ala Phe Trp Leu Ala Leu Leu Pro Ala Leu Phe Leu Leu Ile Gln Ala Ile Gly Ala Pro Phe Gly Tyr Lys Trp Asp Phe Val Ile Leu Asn Gln Gln Leu Ala 35 40 Ala Val Val Asn Ala Ala Phe Ala Leu Leu Ala Ile Val Gly Val Val 55 Ala Asp Pro Thr Thr Ser Gly Leu Gly Asp Ser Asp Arg Val Leu Asn Lys Asp Lys Ser Glu Glu Asn Lys 85 <210> 13 <211> 88 <212> PRT <213> Bacteriophage TPW22 <400> 13 Met Asn Gln Ile Asn Trp Lys Leu Arg Leu Lys Ser Lys Ala Phe Trp 10 Leu Ala Leu Pro Ala Leu Phe Leu Leu Ile Gln Ala Ile Gly Ala

25

Ser Phe Gly Tyr Lys Trp Asn Phe Val Ile Leu Asn Gln Gln Leu Ala Ala Val Val Asn Ala Ala Phe Ala Leu Leu Ala Ile Val Gly Val Val 55 Ala Asp Pro Thr Thr Ser Gly Leu Gly Asp Ser Asp Arg Val Leu Asn Lys Asp Lys Ser Glu Glu Asn Lys 8.5 <210> 14 <211> 74 <212> PRT <213> Artificial Sequence <220> <223> Synthetic Construct <400> 14 Met Arg Phe Asn Met Leu Lys Asn Ser Glu Thr Thr Gly Ala Tyr Val Gly Ser Ala Ile Ala Ile Tyr Ser Gly Phe Thr Leu Ala Asp Trp Ala Ala Ile Phe Gly Ile Leu Phe Gly Leu Phe Thr Met Leu Ile Asn Trp Tyr Tyr Lys Asn Lys Glu Ile Lys Leu Lys Glu Thr Ala Leu Lys Gln Lys Ile Asp Leu Lys Glu Gly Asp His Glu <210> 15 <211> 133 <212> PRT <213> Bacillus phage GA-1 <400> 15

Tyr Phe Leu Leu Gly Ile Ile Gly Val Leu Asn Ile Val Asp Phe Phe 20 25 30

Met Phe Glu Phe Phe His Ser Leu Met Glu Thr Asp Asp Thr Lys Val

Phe Gly Phe Ile Asn Ala Lys Phe Asn Lys Ser Ile Ala Tyr Lys Ser 35 40 45

Ser Lys Thr Ile Asp Gly Ile Met Arg Lys Met Lys Phe Thr Ile Met Ala Ile Leu Phe Ile Pro Val Ser Val Leu Met Pro Glu Pro Ile Gly Leu Gly Ala Leu Tyr Val Phe Tyr Phe Gly Tyr Ile Tyr Ala Glu Leu

Asn Ser Ile Leu Ser His Leu Lys Leu Ser Glu Asp Gly Lys Glu Thr

Glu Val Phe Leu Asp Phe Ile Asn Thr Phe Phe Asn Ser Thr Lys Gly 120

Asp Lys Lys Asp Asp 130

<210> 16 <211> 57 <212> PRT

<213> Staphylococcus phage 187

85

<400> 16

Met Leu Met Val Ile Met Val Gly Asn Val Gly Ile Tyr Leu Thr Ile

Phe Leu Ile Asp Thr Gly Thr Leu Arg His Gln Ala Thr Gln Glu Ile

Trp His Gly Ile Asp Ile Leu Lys Gly Leu Lys Cys Leu Glu Thr Leu

Leu Ile Leu Ser Leu Asn Gln Val Ile 50 55

<210> 17

<400> 17

Met Tyr Gln Met Glu Lys Ile Thr Thr Gly Val Ser Tyr Thr Thr Ser

Ala Val Gly Met Gly Tyr Trp Phe Leu Gln Phe Leu Asp Arg Val Ser

Pro Ser Gln Trp Ala Ala Ile Gly Val Leu Gly Ser Leu Leu Phe Gly Leu Leu Thr Tyr Leu Thr Asn Leu Tyr Phe Lys Ile Arg Glu Asp Arg 55 Arg Lys Ala Ala Arg Gly Glu <210> 18 <211> 75 <212> PRT <213> Artificial Sequence <220> <223> Synthetic Construct <400> 18 Met Glu Arg Trp Thr Leu Leu Asp Ile Leu Ala Phe Leu Leu Leu Leu 10 Ser Leu Leu Pro Ser Leu Leu Ile Met Phe Ile Pro Ser Met Tyr Lys Gln His Ala Ser Leu Trp Lys Ala Arg Ser Leu Ala Lys Thr Leu 40 Ser Met Ala Ser Ser Ala Arg Leu Thr Pro Leu Ser Ser Ser Arg Thr 55 50 Pro Cys Val Leu Lys Gln Asp Ser Lys Lys Leu <210> 19 <211> 87 <212> PRT <213> B.subtilis <400> 19 Met Asn Thr Phe Asp Lys Gly Thr Val Ile Arg Thr Val Leu Leu Ile Ala Leu Ile Asn Gln Thr Met Leu Met Leu Gly Lys Ser Pro Leu

Ala Gly Ser Ile Ala Phe Thr Ile Gly Thr Thr Leu Ala Ala Trp Phe 50 60

Asp Ile Gln Glu Glu Gln Val Asn Gln Leu Ala Asp Ala Leu Tyr Ser 35 40 45

Lys Asn Asn Tyr Val Thr Glu Lys Gly Lys Lys Gln Arg Asp Leu Leu Arg Asp Asn Asn Leu Thr Lys <210> 20 <211> 70 <211> 70 <212> PRT <213> Bacillus subtilis 168 prophage <400> 20 Met Glu Met Asp Ile Thr Gln Tyr Leu Ser Thr Gln Gly Pro Phe Ala Val Leu Phe Cys Trp Leu Leu Phe Tyr Val Met Lys Thr Ser Lys Glu Arg Glu Ser Lys Leu Tyr Asn Gln Ile Asp Ser Gln Asn Glu Val Leu 40 Gly Lys Phe Ser Glu Lys Tyr Asp Val Val Ile Glu Lys Leu Asp Lys Ile Glu Gln Asn Phe Lys <210> 21 <211> 88 <212> PRT <213> Bacillus subtilis 168 prophage <400> 21 Met Phe Glu Asn Ile Asp Lys Gly Thr Ile Val Arg Thr Leu Leu Leu Ala Ile Ala Leu Leu Asn Gln Ile Met Val Met Leu Gly Lys Ala Ala Phe Ile Ile Asn Glu Glu Asp Ile Asn His Leu Tyr Asp Cys Leu Tyr Thr Ile Phe Thr Ile Val Phe Thr Thr Ser Thr Thr Ala Ala Trp Phe Lys Asn Asn Tyr Ile Thr Ala Lys Gly Lys Lys Gln Lys Gln Val

Leu Lys Lys Glu Asn Leu Phe Lys

85

<210> 22

<211> 119 <212> PRT

<213> Bacteriophage phi-Ealh

<400> 22

Met Arg Lys Ile Tyr Val Val Ile Ile Thr Thr Ile Val Val Ala Gly 1 5 10 15

Leu Ile Trp Ala Phe Ile Ala Thr Gln Val Asn Thr Gly Val Thr Ser 20 25 30

Lys Arg Gln Glu Asp Ala Leu Ala Val Ser Glu Ala Asn Val Gly Ile $35 \hspace{1cm} 40 \hspace{1cm} 45$

Gly Lys Glu Ala Lys Asp Gln Gly Glu Gln Ala Thr Lys Arg Ala Asp 50 60

Val Ala Lys Glu Gln Arg Thr His Gln Ile Asn Gln Leu Lys Asp Lys 65 70 75 80

Leu His Glu Lys Ala Glu Ser Tyr Asp Ser Ile Pro Leu Ser Pro Ser 85 90 95

Asp Val Asp Ile Leu Cys Arg Ala Tyr Arg Ser Thr Asp Pro Val Cys 100 105 110

Ser Pro Thr Val Lys Ser Asp 115

<210> 23

<211> 91

<212> PRT

<213> Phage phiX174

<400> 23

Met Val Arg Trp Thr Leu Trp Asp Thr Leu Ala Phe Leu Leu Leu 1 5 10 15

Ser Leu Leu Pro Ser Leu Leu Ile Met Phe Ile Pro Ser Thr Phe 20 25 30

Lys Arg Pro Val Ser Ser Trp Lys Ala Leu Asn Leu Arg Lys Thr Leu 35 40 45

Leu Met Ala Ser Ser Val Arg Leu Lys Pro Leu Asn Cys Ser Arg Leu 50 55 60

Pro Cys Val Tyr Ala Gln Glu Thr Leu Thr Phe Leu Leu Thr Gln Lys Lys Thr Cys Val Lys Asn Tyr Val Gln Lys Glu <210> 24 <211> 109 <212> PRT <213> Artificial Sequence <220> <223> Synthetic Construct <400> 24 Met Pro Cys Leu Ile His Leu Val Gly Trp Gly Ser Ser Pro Gly Ser Ala Leu Ile Arg Glu Gln Ala Ile Gly Ala Gly Leu Ala Ala Trp Met Thr Cys Leu Arg Gly Arg Tyr Leu Gly Arg Gly Trp Arg Lys Thr Thr 40 Phe Asp Ala Ala Ile Cys Ala Leu Ile Ala Trp Phe Ala Arg Asp Gly Leu Ala Leu Val Gly Ile Asp Asn Gln Phe Ser Tyr Leu Ser Ser Ile Ile Val Gly Tyr Leu Gly Asn Asp Tyr Leu Gly Ala Leu Leu Arg Arg 90 Arg Leu Glu Lys Lys Ser Gly Glu Ser Asn Ala Pro Gln <210> 25 <211> 95 <212> PRT <213> Listeria innocua <400> 25 Met Met Lys Met Glu Phe Gly Lys Glu Leu Leu Val Tyr Met Thr Phe Leu Val Val Val Thr Pro Val Phe Val Gln Ala Ile Lys Lys Thr Glu 20

Leu Ile Pro Ser Lys Trp Leu Pro Thr Val Ser Ile Leu Val Gly Ala

Ile Leu Gly Ala Leu Ala Thr Ser Leu Asp Gly Ser Gly Ser Leu Ala

Thr Met Ile Trp Ala Gly Ala Leu Ala Gly Ala Gly Gly Thr Gly Leu

Phe Glu Gln Phe Thr Asn Arg Ala Lys Lys Tyr Gly Lys Asp Asp

<210> 26

<211> 145 <212> PRT <213> Bacteriophage 80 alpha

<400> 26

Met Asp Ile Asn Trp Lys Leu Arg Phe Lys Asn Lys Ala Val Leu Thr 5 10

Gly Leu Val Gly Ala Leu Phe Val Phe Ile Lys Gln Val Thr Asp Leu

Phe Gly Leu Asp Leu Ser Thr Gln Leu Asn Gln Ala Ser Ala Ile Ile

Gly Ala Ile Leu Thr Leu Leu Thr Gly Ile Gly Val Ile Thr Asp Pro 55

Thr Ser Lys Gly Val Ser Asp Ser Ser Ile Ala Gln Thr Tyr Gln Ala

Pro Arg Asp Ser Lys Lys Glu Glu Gln Gln Val Thr Trp Lys Ser Ser

Gln Asp Ser Ser Leu Thr Pro Glu Leu Ser Ala Lys Ala Pro Lys Glu 100 105

Tyr Asp Thr Ser Gln Pro Phe Thr Asp Ala Ser Asn Asp Val Gly Phe 115

Asp Val Asn Glu Tyr His His Gly Gly Gly Asp Asn Ala Ser Lys Ile

Asn 145

<210> 27

<211> 145 <212> PRT

<213> Staphylococcus bacteriophage phi 11

<400> 27

Met Asp Ile Asn Trp Lys Leu Arg Phe Lys Asn Lys Ala Val Leu Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Gly Leu Val Gly Ala Leu Phe Val Phe Ile Lys Gln Val Thr Asp Leu 20 25 30

Phe Gly Leu Asp Leu Ser Thr Gln Leu Asn Gln Ala Ser Ala Ile Ile 35 40 45

Thr Ser Lys Gly Val Ser Asp Ser Ser Ile Ala Gln Thr Tyr Gln Ala 65 70 75 80

Pro Arg Asp Ser Lys Lys Glu Glu Gln Gln Val Thr Trp Lys Ser Ser 85 90 95

Gln Asp Ser Ser Leu Thr Pro Glu Leu Ser Ala Lys Ala Pro Lys Glu 100 105 110

Tyr Asp Thr Ser Gln Pro Phe Thr Asp Ala Ser Asn Asp Val Gly Phe 115 120 125

Asp Val Asn Glu Tyr His His Gly Gly Gly Asp Asn Ala Ser Lys Ile 130 135 140

Asn 145

<210> 28

<211> 138

<212> PRT

<213> Streptococcus pneumoniae bacteriophage MM1

<400> 28

Met Lys Ile Glu Phe Phe Asn Phe Leu Arg Ser Val Ile Gln Thr Glu $1 ag{5} ag{5}$

Asp Gly Leu Val Leu Tyr Ala Leu Ala Leu Ile Val Ser Met Glu Ile 20 25 30

Ile Asp Phe Val Thr Gly Thr Ile Ala Ala Ile Ile Asn Pro Asp Ile 35 40 45

Glu Tyr Lys Ser Lys Ile Gly Ile Asn Gly Leu Leu Arg Lys Ile Ser

60

Gly Val Leu Leu Met Ile Leu Ile Pro Ala Ser Val Leu Leu Pro 70 Glu Lys Thr Gly Phe Ala Phe Leu Tyr Ser Ile Cys Leu Gly Tyr Ile Ala Phe Thr Phe Gln Ser Leu Ile Glu Asn Tyr Arg Lys Leu Lys Gly Asn Val Thr Leu Phe Gln Pro Ile Val Lys Val Phe Gln Arg Leu Leu 120 115 Glu Lys Asp Asp Asp Thr Lys Lys Gly Glu <210> 29 <211> 86 <212> PRT <213> Streptococcus thermophilus bacteriophage Sfi21 <400> 29 Met Lys Lys Arg Lys Lys Met Ile Asn Phe Lys Leu Arg Leu Gln Asn Lys Ala Thr Leu Val Ala Leu Ile Ser Ala Val Phe Leu Met Leu · 20 25 Gln Gln Phe Gly Leu His Val Pro Asn Asn Ile Gln Gly Ile Asn Thr Leu Val Gly Ile Leu Val Ile Leu Gly Ile Ile Thr Asp Pro Thr Thr Lys Gly Ile Ala Asp Ser Glu Arg Ala Leu Ser Tyr Ile Gln Pro Leu 70 Asp Asp Lys Glu Val Tyr <210> 30 <211> 96 <212> PRT <213> Bacteriophage A500 <400> 30 Met Met Lys Met Glu Phe Gly Lys Glu Leu Leu Val Tyr Met Thr Phe 1.0

55

50

Leu Val Val Val Thr Pro Val Phe Val Gln Ala Ile Lys Lys Thr Glu

Leu Ile Pro Ser Lys Trp Leu Pro Thr Val Ser Ile Leu Val Gly Ala

Ile Leu Gly Ala Leu Ala Thr Ser Leu Asp Gly Ser Gly Ser Leu Ala 55

Thr Met Ile Trp Ala Gly Ala Leu Ala Gly Ala Gly Gly Thr Gly Leu 70

Phe Glu Gln Phe Thr Asn Arg Ala Lys Lys Tyr Gly Lys Asp Asp Lys

<210> 31

<211> 90 <212> PRT

<213> Bacteriophage PL-1

<400> 31

Met Gln Asn Glu Leu Leu Gln Val Leu Ala Ile Ala Phe Val Ile Ala

Pro Ile Thr Thr Gly Phe Thr Glu Ile Phe Lys Arg Tyr Thr Pro Ala

Glu Gly Lys Leu Leu Pro Val Leu Ser Ile Gly Thr Gly Ile Leu Leu

Ala Cys Val Trp Ala Met Ala Phe Gly His Leu Pro Leu Ile Gly Ala

Tyr Ala Leu Ala Gly Met Leu Ser Gly Leu Ala Ser Val Gly Val Tyr

Gln Ile Val Lys Pro Asn Glu Glu Val Lys 85